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APR ● ﹍﹍
1643
TECH CENTER 1600/2900

N / A

#8

```
This Raw Listing contains the General Information
Section and up to first 5 pages.
```

ENTERED R.S

```
 1   <110> APPLICANT: GILLIES, Stephen D
 2         LO, Kin-Ming
 3         LAN, Yan
 4         WESOLOWSKI, John
 5   <120> TITLE OF INVENTION: Enhancing the Circulating Half-life of Antibody-based
 6         Fusion Proteins
 7   <130> FILE REFERENCE: LEX-003
 8   <140> CURRENT APPLICATION NUMBER: US/09/256,156A
 9   <141> CURRENT FILING DATE: 1999-02-24
10   <150> EARLIER APPLICATION NUMBER: US 60/075,887
11   <151> EARLIER FILING DATE: 1998-02-25
12   <160> NUMBER OF SEQ ID NOS: 8
13   <170> SOFTWARE: PatentIn Ver. 2.0
14   <210> SEQ ID NO 1
15   <211> LENGTH: 447
16   <212> TYPE: PRT
17   <213> ORGANISM: Homo sapiens
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19   <223> OTHER INFORMATION: IGG-1 CHAIN C REGION
20   <220> FEATURE:
21   <221> NAME/KEY: VARIANT
22   <222> LOCATION: (1)..(117)
23   <223> OTHER INFORMATION: The Xaa at positions 1 to 117 are non-conserved
24         amino acids
25   <400> SEQUENCE: 1
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      31                  35                  40                  45
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      33              50                  55                  60
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      39                  100                 105                 110
W-->  40      Xaa Xaa Xaa Xaa Xaa Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu
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      43              130                 135                 140
      44      Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
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1643

APR 13 20.

PAGE: 2   **RAW SEQUENCE LISTING**   DATE: 06/30/1999
                     PATENT APPLICATION US/09/256,156A   TIME: 16:57:39

Input Set: **I256156A.RAW**

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49              180                185                190
50          Leu Gly Thr Gln Thr Tyr Ile Cys Asn Val Asn His Lys Pro Ser Asn
51              195                200                205
52          Thr Lys Val Asp Lys Lys Val Glu Pro Lys Ser Cys Asp Lys Thr His
53              210                215                220
54          Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val
55          225                230                235                240
56          Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr
57              245                250                255
58          Pro Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu
59              260                265                270
60          Val Lys Phe Asn Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys
61              275                280                285
62          Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser
63              290                295                300
64          Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys
65          305                310                315                320
66          Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile
67              325                330                335
68          Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro
69              340                345                350
70          Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu
71              355                360                365
72          Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn
73          370                375                380
74          Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser
75          385                390                395                400
76          Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg
77              405                410                415
78          Trp Gln Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu
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83   <211> LENGTH: 443
84   <212> TYPE: PRT
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91   <223> OTHER INFORMATION: The Xaa at positions 1 to 117 are non-conserved
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W--> OK

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W-->        98    Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
            99                35                40                45
W-->       100    Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
           101           50                55                60
W->        102    Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
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W-->       104    Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
           105                85                90                95
W-->       106    Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
           107                100               105               110
W-->       108    Xaa Xaa Xaa Xaa Xaa Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu
           109                115                120               125
           110    Ala Pro Cys Ser Arg Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys
           111           130                135               140
           112    Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
           113        145                150               155               160
           114    Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser
           115                165               170               175
           116    Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Asn
           117                180               185               190
           118    Phe Gly Thr Gln Thr Tyr Thr Cys Asn Val Asp His Lys Pro Ser Asn
           119                195               200               205
           120    Thr Lys Val Asp Lys Thr Val Glu Arg Lys Cys Cys Val Glu Cys Pro
           121           210               215               220
           122    Pro Cys Pro Ala Pro Pro Val Ala Gly Pro Ser Val Phe Leu Phe Pro
           123        225                230               235               240
           124    Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr
           125                245               250               255
           126    Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val Gln Phe Asn
           127                260               265               270
           128    Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr Lys Pro Arg
           129                275               280               285
           130    Glu Glu Gln Phe Asn Ser Thr Phe Arg Val Val Ser Val Leu Thr Val
           131           290               295               300
           132    Val His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser
           133        305               310               315               320
           134    Asn Lys Gly Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser Lys Thr Lys
           135                325               330               335
           136    Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser Arg Glu
           137                340               345               350
           138    Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe
           139                355               360               365
           140    Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Asn Gly Gln Pro Glu
           141        370               375               380
           142    Asn Asn Tyr Lys Thr Thr Pro Pro Met Leu Asp Ser Asp Gly Ser Phe
           143        385               390               395               400
           144    Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly
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145                              405                 410                 415
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159     <223> OTHER INFORMATION: The Xaa at positions 1 to 117 are non-conserved
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        165   20   25   30
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        167   35   40   45
W-->   168   Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        169   50   55   60
W-->   170   Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
        171   65   70   75   80
W-->   172   Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
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W-->   174   Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
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W-->   176   Xaa Xaa Xaa Xaa Xaa Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu
        177   115   120   125
        178   Ala Pro Cys Ser Arg Ser Thr Ser Gly Gly Thr Ala Ala Leu Gly Cys
        179   130   135   140
        180   Leu Val Lys Asp Tyr Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser
        181   145   150   155   160
        182   Gly Ala Leu Thr Ser Gly Val His Thr Phe Pro Ala Val Leu Gln Ser
        183   165   170   175
        184   Ser Gly Leu Tyr Ser Leu Ser Ser Val Val Thr Val Pro Ser Ser Ser
        185   180   185   190
        186   Leu Gly Thr Gln Thr Tyr Thr Cys Asn Val Asn His Lys Pro Ser Asn
        187   195   200   205
        188   Thr Lys Val Asp Lys Arg Val Glu Leu Lys Thr Pro Leu Gly Asp Thr
        189   210   215   220
        190   Thr His Thr Cys Pro Arg Cys Pro Glu Pro Lys Ser Cys Asp Thr Pro
        191   225   230   235   240
        192   Pro Pro Cys Pro Arg Cys Pro Glu Pro Lys Ser Cys Asp Thr Pro Pro
        193   245   250   255
        194   Pro Cys Pro Arg Cys Pro Glu Pro Lys Ser Cys Asp Thr Pro Pro Pro

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      195                    260                  265                  270
      196          Cys Pro Arg Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe
      197                    275                  280                  285
      198          Leu Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro
      199                    290                  295                  300
      200          Glu Val Thr Cys Val Val Val Asp Val Ser His Glu Asp Pro Glu Val
      201          305                  310                  315                  320
      202          Gln Phe Lys Trp Tyr Val Asp Gly Val Glu Val His Asn Ala Lys Thr
      203                    325                  330                  335
      204          Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Phe Arg Val Val Ser Val
      205                    340                  345                  350
      206          Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Lys Cys
      207                    355                  360                  365
      208          Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr Ile Ser
      209                    370                  375                  380
      210          Lys Thr Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro
      211          385                  390                  395                  400
      212          Ser Arg Glu Glu Met Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val
      213                    405                  410                  415
      214          Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser Ser Gly
      215                    420                  425                  430
      216          Gln Pro Glu Asn Asn Tyr Asn Thr Thr Pro Pro Met Leu Asp Ser Asp
      217                    435                  440                  445
      218          Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp
      219                    450                  455                  460
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      234       amino acids
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      239                    20                  25                  30
W-->  240          Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      241                    35                  40                  45
W-->  242          Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
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W-->  244          Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing t ensure that a corresponding explanation is presented in the <220> to
<223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION **US/09/256,156A**

Input Set: **I256156A.RAW**

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26	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
28	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
30	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
32	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
34	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
36	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
38	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
40	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Ala Ser Thr Lys Gly P
94	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
96	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
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162	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
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236	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa X
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250	W	"N" or "Xaa" used: Feature required	Xaa Xaa Xaa Xaa Xaa Ala Ser Thr Lys Gly P